

EMBOSS_001	177	ccttcataatcggtgccagaggtgaagaactggccaatgttaaggcg	226
EMBOSS_001	243	CTTCATCGATGCGGTGCCATCATGCTGGATGTGGCCAATGTTGCCGGG	292
EMBOSS_001	227	agcagttctccaacatggccagcgagacaatgacccggtgacgtggtgctt	276
EMBOSS_001	293	TCAGGTGGCCAAGTGGGAGCAGGATATCACTCTGACATCTGTATT	342
EMBOSS_001	277	aagctgtccagcgggtgaacgagc-----tgctagcacgggacgatgt	320
EMBOSS_001	343	TCCATGTGTCAGAAGCTGAAC-CGGTGTGATGTGAGGACCCGACCA--T	389
EMBOSS_001	321	ggacggtgttgtcatcaccacgggcaccgacacccgtggaggagtctgcct	370
EMBOSS_001	390	GGCCGGTGTGTTCATACCCACGGCACCGACCCCTCGAGGAGACTGCCT	439
EMBOSS_001	371	acttcctgcacctcactgtcaaaagtgcacaagccagttgtcttgtggct	420
EMBOSS_001	440	TCCTTCGGACGCCACTGTGCAACTGTGGCAAGCCAATTGTTCATCTGGGT	489
EMBOSS_001	421	gccatgcgccagccacggccatctcagctgacggggcccatgaatctgct	470
EMBOSS_001	490	GCCATGCGGCCATCCACGGCCATCTCAGCTGACGGGCCCTCAACTCTGT	539
EMBOSS_001	471	cgaagccgtgcgggtggctggcgacaagcagtcgcgcggtgcgggtgtca	520
EMBOSS_001	540	CGAAGCCGTGACGGTGGCTGCCTCACAGTCGCGCGCGGATCGCGGTGCCA	589
EMBOSS_001	521	tggtggtctcacaagcatgcattggctcgggccgctatatcaccaagacc	570
EMBOSS_001	590	TGTTGGTCATGAACGATCGCATTGCTCGGCCCTACTATGTACCAAGACC	639
EMBOSS_001	571	aatgcctccactctggacaccttcaaggccaatgaggagggtcactctgg	620
EMBOSS_001	640	AATGCCAACACTATGGACACCTTCAAGGCCATGGAGATGGGCTACCTTGG	689
EMBOSS_001	621	cgtgatcatcgggcaac-cgcatt-----tactaccaaaccgatatcgaca	664
EMBOSS_001	690	CGAGATGATCTCAACACCCCTTCTTCTTCTAC-----CGG-CCGTC	733
EMBOSS_001	665	agctgcataccactcgggt-----cggtctttgacgtccgcggcctgacttc	710
EMBOSS_001	734	AG-----CCAACGGTAAGGTGGCCTTTGACATCACCAACGTGACTGA	776
EMBOSS_001	711	gctccccaagtggacattcttgtaggttatcaggac---gaccccgaa	757
EMBOSS_001	777	GATCCCCGTTGGACATCTGTTTTCTTATGAGGACATGCACACGACA	826
EMBOSS_001	758	acctctacgacgcgcgcacatcagcatggtgtcaagggaattgtgtatgcc	807
EMBOSS_001	827	CCCTCTACAACGCCATC-TCCAG--TGGTGCCAGGGAATTGTGATTGCC	873
EMBOSS_001	808	gggatgggtgctggatcgctctca-----gttc-----gtgg-tat-----	842
EMBOSS_001	874	GGGGCTGTGTCTGGAGGCGGTCAACCTCCTTCAATGAGGCTATCGAGGA	923
EMBOSS_001	843	tgcggcgatcgtaaggctatggagaagggtgctgtgtcatcaggagta	892
EMBOSS_001	924	TGTCATCAACCG-----TTTGAGA-----TCCTGTCTGTGAGAGTA	961
EMBOSS_001	893	cgcgacaggcaatgggatagtgccac---ccgac--gaagagctccccc	937
EMBOSS_001	962	TGCGCACAGTCAATGGGGAAGTGCCACTGTGACAGCTGACAGCGACACC	1011
EMBOSS_001	938	gcc-----tcgtcagtgactccctaaaccggcgcatgccccgattct	980
EMBOSS_001	1012	GCCACCCCATCGCCAGTGGAATACCTAAACCCGAGAGTCCCGCATTCT	1061
EMBOSS_001	981	gttgatgttggcgctaactc--ggacaag--tgatcctaagtaatccaa	1026

|| ||.|||.| ||.||||.
SEQ ID NO:2 EMBOSS_001 1108 GACGTGTTTGCTCTGGGCACGGATGCGTAG 1137

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